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AT BE CH DE FR GB IT LI LU NL SE(71) Applicant: CHIRON CORPORATION  
4560 Horton Street  
Emeryville California 94608(US)(71) Applicant: The Board of Trustees of the University of  
Washington  
Seattle Washington(US)(72) Inventor: Agabian, Nina  
3 Burnett Avenue N., No. 5  
San Francisco California 94131(US)(72) Inventor: Stephens, Richard  
2715 W. Blaine  
Seattle Washington 98199(US)(72) Inventor: Kuo, Cho-Chou  
6531 29th Avenue N.E. Seattle  
Washington 98115(US)(72) Inventor: Mullenbach, Guy T.  
309 63rd Street No. C, Oakland  
California 94618(US)(74) Representative: Glawe, Delfs, Moll & Partner  
Patentanwälte  
Postfach 162 Liebherrstrasse 20  
D-8000 München 26(DE)(54) *Chlamydia* major outer membrane protein.

(57) Methods and compositions are provided for the production of a polypeptide which is immunologically cross-reactive with a naturally-occurring major outer membrane protein (MOMP) of *Chlamydia trachomatis*. A DNA construct including a replication system recognized by *E. coli*, and an MOMP gene under the transcriptional control of a  $\beta$ -galactosidase promoter and terminator is provided.

Recombinant phage  $\lambda$ gt11/L2/33 was deposited at the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852, on January 10, 1985 and granted accession no. 40157. L2 B9-F DNA was deposited at the American Type Culture Collection on December 31, 1985, and granted accession no 40217.

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polynucleotide fragment capable of binding a DNA or RNA  
sequence characteristic of one or more species of  
Chlamydia trachomatis, and a detectable label bound to  
the single-stranded polynucleotide.

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art of

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25. A polynucleotide probe as in claim 24,  
wherein the sequence of the single stranded  
polynucleotide fragment is substantially homologous or  
complementary to at least 12 contiguous bases as set  
forth in Appendix B hereto.

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various MOMP's generally ranges from about 38kD to 45kD. The serovars display varying antigenic complexity, with certain serovars eliciting broad cross-reactivity with others in the same biovar, while  
5 other serovars display little or no such cross-reactivity.

Vaccines utilizing purified and unpurified preparations of intact Chlamydia trachomatis have been prepared and tested on monkeys. While successful  
10 protection against subsequent challenge with the same chlamydial serovar was achieved, it was found that heterologous serovar challenge resulted in more severe pathology than that experienced by controls who had not been immunized. In human trials, immunization with the  
15 vaccines afforded significant protection against the serovar of the vaccine for up to two years, but hypersensitivity resulted from infection with heterologous serovars.

## 2. Description of the Relevant Art

20 The nature of the major outer membrane protein and its relation to the biovars and serovars of Chlamydia trachomatis are discussed in Grayston and Wang (1975) J. Infect. Dis. 132:87-105; Stephens et al. (1982) J. Immunol. 128:1083-1089; and Caldwell et al.  
25 (1981) Infect. Immun. 31:1161-1176. Inhibition of infectivity of Chlamydia trachomatis by both anti-chlamydial antisera and monoclonal antibodies has been demonstrated. Caldwell and Perry (1982) Infect. Immun. 38:745-754; and Clark et al. (1982) Infect.  
30 Immun. 38:1273-1278. Vaccine trials conducted with intact chlamydial elementary bodies are reported by Collier (1961) Lancet 1:795-800; Wang et al. (1967) Amer. J. Ophthal. 63:1615-1630; and Woolridge et al. (1967) Amer. J. Ophthal. 63:1645-1653. The cloning and  
35 expression of a gene encoding a 74,000 dalton chlamydial antigen in E. coli is reported by Stephens et al. (1983) Abstracts Annual Meeting American Society

of Microbiology, B29, p. 35. Stephens et al. failed to obtain expression of a major outer membrane protein. Wenman and Lovett (1982) Nature 296:68-70, report the expression of a 19,000 dalton Chlamydia trachomatis polypeptide. The polypeptide does not appear to be involved in the major outer membrane protein. Allan et al. (1984) Infect. Immun. 45:637-641, recently reported the cloning of the major outer membrane protein gene. Nano et al. (1985) Infect. Immun. 45:637-641 report the sequencing of the first 25 N-terminal amino acids of the major outer membrane protein and the cloning of at least a portion of the gene. An immunoassay for the detection of Chlamydia trachomatis antigen is described in U.S. Patent No. 4,497,899.

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#### SUMMARY OF THE INVENTION

Polypeptide compositions having immunological activity corresponding to that of a major outer membrane protein (MOMP) of Chlamydia trachomatis are produced by expressing a chimeric DNA construct comprising a polynucleotide encoding at least a portion of the MOMP under the regulatory control of a regulatory system recognized by a unicellular expression host. The MOMP polynucleotide may code for the entire protein or for a fragment thereof, and may be expressed in conjunction with another structural gene to yield a fused translation product. Such polypeptide compositions will be characterized by the presence of non-interfering amounts of substances derived from the expression, which presence may be used to distinguish the polypeptides of the present invention from the natural polypeptides. The polypeptide compositions of the present invention are useful as substitutes for the naturally-occurring MOMP's of Chlamydia trachomatis, particularly as immunological reagents, e.g., in serological assays to detect the presence of antibodies in blood, the immunogenic substance in vaccines, and the like. The

MOMP polynucleotides will also be useful as labelled probes for diagnostics.

#### DESCRIPTION OF THE SPECIFIC EMBODIMENTS

Methods and compositions are provided for the efficient expression of polypeptides demonstrating immunological activity analogous to that of a major outer membrane protein (MOMP) of Chlamydia trachomatis. By analogous immunological activity, it is meant that, when administered to a vertebrate, the polypeptides will elicit an immunological response which is cross-reactive with antibodies elicited by administration of the natural MOMP. The present invention employs a DNA construct including a MOMP polynucleotide or fragment thereof encoding a polypeptide under the transcriptional and translational control of suitable regulatory sequences. MOMP polypeptides free from other chlamydial antigens are obtained by expressing the MOMP polynucleotides in a unicellular host other than Chlamydia trachomatis.

The MOMP polynucleotides of the present invention may be initially derived from any of the chlamydial serovars and may be employed in a natural or modified form. MOMP nucleotides having a sequence corresponding to an intact natural MOMP gene will usually be employed when it is desired to express the entire MOMP, although it will be possible to alter the sequence for a desired purpose, e.g., to conform to the codon bias of the expression host (as discussed below), or to create or delete restriction sites, so long as the amino acid sequence is not significantly altered. Shorter DNA sequences corresponding to portions of the MOMP gene will be employed when it is desired to produce only a fragment of the natural proteins. Usually, such sequences will encode for an epitopic region(s), comprising at least 27 bp, usually at least 36 bp, preferably at least 45 bp, and may be much longer.

It will sometimes be desirable to express the MOMP polynucleotide together with other gene(s) in order to provide fused translation products having desirable properties. For example, when producing low molecular weight MOMP polypeptide fragments (below about 5kD), it may be desirable to fuse the MOMP polypeptide to an immunogenic carrier, e.g., tetanus toxoid or hepatitis B surface antigen. It is also possible to fuse MOMP fragments from more than one serovar to each other and/or a gene expressing a suitable immunogenic carrier. Conveniently, the MOMP polynucleotide will be inserted in proper reading frame with the fused gene and under the regulatory control of the regulatory system of the fused gene. Recovery of the gene product may be facilitated by employing a secretory protein as the fused gene product, as described below.

The MOMP polypeptides may be glycosylated, partially glycosylated, or unglycosylated, depending on the nature of the expression host. Generally, prokaryotes such as E. coli will provide no glycosylation of the translated MOMP gene products, while yeast and mammalian cell culture will provide partial or substantial glycosylation. Thus, it will be possible to vary the final MOMP product by appropriate selection of the expression host.

The MOMP polynucleotide may be synthetic or natural, or combinations thereof. A natural MOMP gene (or a portion thereof) may be obtained by preparing a Chlamydia trachomatis genomic library and screening for the presence of the MOMP gene. Screening may be accomplished using antibodies for the gene product or using labelled DNA probes specific for the polynucleotide. Both methods are exemplified in the Experimental section hereinafter. Suitable antibodies are commercially available or may be prepared from purified MOMP obtained from Chlamydia trachomatis by

well known techniques. Suitable DNA probes may be obtained based on the amino acid sequence of the MOMP, or based on the polynucleotide sequence which is reported hereinafter for the MOMP of the L<sub>2</sub> serovar (see Appendix B). Conveniently, the  $\lambda$ gt11/L2/33 clone which has been deposited in connection with this patent application may be labelled and used as a screening probe. A specific method for selecting a clone expressing the MOMP of the L<sub>2</sub> serovar from a Chlamydia trachomatis genomic library is set forth in the Experimental section hereinafter. This method can be modified to allow for selection of MOMP gene(s) from other serovars.

Synthetic polynucleotide sequences encoding for at least a portion of the MOMP gene of Chlamydia trachomatis may also find use, either alone or in combination with the naturally-occurring sequences. Coding for the synthetic sequences may be based on either the reported amino acid sequences for the MOMP's or on the polynucleotide sequences which are determined from the MOMP genes by known techniques. When used for preparing polypeptides as immunological reagents or as vaccines, it is usually desirable that the synthetic nucleotide fragment code for an oligopeptide corresponding to an epitopic site of the natural MOMP. Often, such epitopic sites may be inferred from the folding rules of Chou and Fasman (1974) Biochemistry 13:211-222, in conjunction with an analysis of hydrophobic and hydrophilic regions of the protein as taught by Hopp and Woods (1981) Proc. Natl. Acad. Sci. USA 78:3824-3828. Alternatively, the DNA sequences encoding the polypeptide regions which react with particular monoclonal antibodies may be identified by the high-density phage procedure described by Nunberg et al. (1984) Proc. Natl. Acad. Sci. USA 81:3675-3679. The oligopeptide may then be screened for eliciting Ab cross-reactive with the naturally-occurring MOMP.

A number of techniques are available for synthesizing short, single-stranded DNA fragments, e.g., the phosphoramidite method described by Beaucage and Carruthers (1981) Tetrahedron Lett. 22:1859-1862.

5 A particularly useful adaptation of the method of Beaucage and Carruthers is reported by Warner et al. (1984) DNA 3:401-411. Using the method of Urdea et al., single-stranded DNA fragments having a length of up to 100 bases may be synthesized, and double-stranded  
10 DNA fragments may be formed by annealing and ligating a plurality of single-stranded fragments under appropriate conditions. Alternatively, the complementary strand may be added using DNA polymerase with an appropriate primer sequence.

15 When preparing synthetic MOMP polynucleotides, it may sometimes be desirable to modify the natural nucleotide sequence. For example, it will often be preferred to use codons which are preferentially recognized by the desired host. When  
20 employing a yeast host, codons which appear at high frequency in the structural genes encoding the yeast glycolytic enzymes may be employed. In some instances, it may be desirable to further alter the nucleotide sequence to create or remove restriction sites to  
25 increase stability or to substitute one or more amino acids in the resulting polypeptide. Such changes may be made to enhance the immunogenicity of the polypeptide, facilitate conjugating the polypeptide to a carrier protein, or the like. It may also be  
30 desirable to add amino acids to the N-terminus or C-terminus of the polypeptide, where such additional amino acids provide for a desired result.

To produce a desired MOMP polypeptide, the MOMP polynucleotides will be incorporated into DNA  
35 constructs capable of being introduced into a desired expression host, usually either a prokaryotic host or eukaryotic host, such as yeast. Such DNA constructs



will include the MOMP polynucleotide encoding the polypeptide product, transcriptional and translational initiation regulatory sequences joined to the 5'-end of the polynucleotide, and transcriptional and translational termination regulatory sequences joined to the 3'-end of the polynucleotide. The DNA constructs will usually also include a replication system recognized by the expression host to allow for self-replication, and will often include other functional sequences such as markers allowing for selection of transformed hosts, additional replication systems, secretory leader and processing signal sequences, and the like. The replication system, however, is not necessary since the DNA construct may allow for integration into the host genome. Integration is facilitated by providing short DNA fragments on either side of the MOMP polynucleotide, which fragments are homologous to a desired location in the host genome.

The transcriptional initiation regulatory sequences will include a promoter region recognized by the expression host. For E. coli hosts, the lac promoter, lambda P<sub>L</sub> or P<sub>R</sub>, or the  $\beta$ -galactosidase promoter, as exemplified in the Experimental section hereinafter, are suitable. For yeast hosts, suitable promoters include those involved with the enzymes in a yeast glycolytic pathway, such as the promoters for alcohol dehydrogenase, glyceraldehyde-3-phosphate dehydrogenase, pyruvate kinase, triose phosphate isomerase, phosphoglucose isomerase, phosphofructokinase, and the like. By employing these promoters with other regulatory sequences, such as enhancers, operators, and the like, and using a host having an intact regulatory system, one can regulate the expression of the MOMP polypeptide by a number of techniques, such as varying the carbon source, e.g., replacing glucose with galactose; varying the concentration of a nutrient,

e.g., acid phosphatase, or changing the temperature with a temperature sensitive promoter or regulatory system.

5       The transcriptional termination regulatory  
sequence will include a terminator, preferably a  
terminator balanced with the promoter to provide proper  
transcription. Conveniently, the terminator which is  
naturally found with the promoter may be employed. In  
the exemplary embodiment described in the Experimental  
10   section hereinafter, the MOMP polynucleotide is  
inserted between the  $\beta$ -galactosidase promoter and  
terminator within the  $\beta$ -galactosidase structural gene  
so that a fusion product is formed.

Enhanced yields of the polypeptides of the  
15   present invention may be obtained by employing DNA  
constructs which include a secretory leader and  
processing signal sequence to effect secretion of the  
gene product in yeast. The use of such secretory  
leader and processing signal sequences will be  
20   particularly effective with polypeptides below about 40  
kilodaltons, more usually below about 30 kilodaltons,  
although it is expected that the system will function  
with polypeptides equal to the length of the whole  
MOMP, i.e., ranging from 38 to 45 kilodaltons. The  
25   secretory leader and processing signal sequences will  
normally be derived from naturally-occurring DNA  
sequences in yeast which provide for secretion of a  
polypeptide. Such polypeptides which are naturally  
secreted by yeast include  $\alpha$ -factor,  $\alpha$ -factor, acid  
30   phosphatase, and the like. If desired, the  
naturally-occurring sequence may be modified, for  
example, by reducing the number of lys-arg pairs in  
 $\alpha$ -factor which define the processing site (while  
retaining at least one pair), or by reducing the length  
35   of the secretory leader sequence (while retaining  
sufficient length to provide for secretion) or by  
introducing point mutations, deletions or other

modifications which facilitate manipulation, e.g.,  
introducing restriction recognition sites.

Conveniently, the secretory leader and processing  
signal sequence may be joined to the MOMP

5 polynucleotide by providing appropriate cohesive ends  
on the polynucleotide fragment, by use of appropriate  
adaptor molecules, or a combination of both. A portion  
of the structural gene for the secretory protein may be  
left in the final DNA construct when it is desired to  
10 produce a fused translation product, as discussed  
above.

Polypeptides of the present invention may  
also be recovered intracellularly as follows. After  
the transformed cell culture has reached a high  
15 density, the cells will be separated, typically by  
centrifugation, lysed, and the MOMP polypeptides  
isolated and purified by various techniques, such as  
extraction, affinity chromatography, electrophoresis,  
dialysis, and combinations thereof.

20 The MOMP polypeptides may also be prepared by  
conventional solid-phase synthesis techniques, such as  
those described by Merrifield (1963) J. Am. Chem. Soc.  
85:2149-2156. Such solid-phase techniques are suitable  
for preparation of polypeptide fragments of up to about  
25 50 to 100 amino acids, or more. Generally, however, as  
the length of the polypeptide increases above 25 amino  
acids, the difficulty in the synthesis increases and  
the desirability of employing a solid-phase synthesis  
technique diminishes.

30 The polypeptides of the present invention,  
and fragments thereof, may be employed in a variety of  
ways. The polypeptides can be employed both as  
labelled and unlabelled reagents in various  
immunoassays, bioassays, and the like, for the  
35 detection of Chlamydia trachomatis or antibodies to  
Chlamydia trachomatis in a biological sample, e.g.,  
serum. Suitable labels include radionuclides,

enzymes, fluorescers, chemiluminescers, enzyme substrates or co-factors, enzyme inhibitors, particles, dyes, and the like. Such labelled reagents may be used in a variety of well known assays, such as

5 radioimmunoassays, enzyme immunoassays, e.g., ELISA, fluorescent immunoassays, and the like. See, for example, U.S. Patent Nos. 3,766,162; 3,791,932; 3,817,837; 3,996,345; and 4,233,402. Polypeptides of the present invention may also find use in vaccines

10 against infection by Chlamydia trachomatis. Larger polypeptides, having a molecular weight exceeding about 5,000 daltons, may be used without further modification. Smaller haptens (i.e., those below about

15 immunogenic carrier in order to elicit the desired immune response. Suitable immunogenic carriers include tetanus toxoid and hepatitis B surface antigen. It will be possible to link short DNA fragments expressing the MOMP polypeptides to genes expressing proteins from

20 other pathogenic organisms or viruses. In this way, the resulting fused proteins may provide immunity against more than one disease.

In preparing a vaccine, the polypeptides will normally be incorporated in a physiologically

25 acceptable medium, such as water, physiological saline, phosphate buffered saline, and the like. The vaccine may be administered intravenously, intraarterially, subcutaneously, intraperitoneally, or the like. The amount of immunogen employed per dose will be about 5

30 to 10 micrograms, if liquid, in a volume of about 0.25 to 1 ml, and may be administered repeatedly at about 2 to 4 week intervals, usually not more than 2 or 3 times.

The polynucleotides of the present invention

35 may be employed as labelled polynucleotide probes suitable for screening biological samples for the presence of various strains of Chlamydia trachomatis.

Probes comprising DNA or RNA from conserved regions of the MOMP gene may be employed for detecting a broad range of Chlamydia, while probes comprising regions of the MOMP gene specific for a particular strain may be employed to identify that strain. The polynucleotide sequences in such probes will typically be at least 12 nucleotides, more typically 16 or more nucleotides. Conveniently, the nucleotide fragment may be synthesized based on the sequence set forth in Appendix A, hereinafter.

Suitable labels include radionuclides, heavy metals, organic ligands, and the like, which allow for detection in conventional assays. Biological samples will be prepared in a conventional manner, e.g., by lysing the Chlamydia to release the nucleic acids.

#### EXPERIMENTAL

The following experiments are offered by way of illustration, not by way of limitation.

#### MATERIALS AND METHODS

##### Reagents

DNase, RNase, endonuclease restriction enzymes, T4 ligase, kinase, DNA Polymerase I, and EcoRI methylase were obtained from Bethesda Research Laboratories. Nitrocellulose was obtained from Schleicher and Schuell. Peroxidase conjugated anti-mouse, anti-rabbit, and peroxidase anti-peroxidase (PAP) sera were obtained from Cappel. Proteinase K, isopropylthiogalactoside (IPTG), and 4-chloro-1-naphthol were from Sigma Chemical Co. Phage packaging mix was obtained from Amersham. CNBr-activated Sepharose<sup>®</sup>-4B was obtained from Pharmacia.

##### Bacterial Strains

E. coli Y1088, Y1089, Y1090, and BNN 97 were obtained from R. Young and R. Davis (Stanford University). For C. trachomatis, two trachoma strains, B/TW-5/OT, and C/TW-3/OT, and one LGV strain,

L<sub>2</sub>/434/Bu, were grown in HeLa 229 cells and Renografin purified as described by Kuo et al. (1977) in:

"Nongonococcal Urethritis and Related Infections,"  
Hobson and Holmes, eds., Am. Soc. Microbiol. pp.  
5 176-185.

#### Antibodies

Polyvalent antiserum to C. trachomatis was obtained from rabbits immunized with purified LVG (L<sub>2</sub> serovar) organisms that were grown in chick embryo yolk  
10 sacs. Anti-E. coli reactivities in this antiserum were removed by passage through a Sepharose<sup>®</sup>-4B column derivitized with an E. coli lysate. For this purpose, approximately 20mg of DNase and RNase treated lysate of induced BNN 97 were coupled to 1mg of CNBr-activated  
15 Sepharose<sup>®</sup>-4B according to the manufacturer's instructions. The development, specificities, and ascites production of monoclonal antibodies specific for C. trachomatis have been previously reported (Stephens et al. (1982) J. Immunol. 128:1083).

#### 20 Insertion of Chlamydial DNA into $\lambda$ gt11

Chlamydial DNA was isolated from cell extracts of serovars L<sub>2</sub>, B, and C by proteinase K treatment (65 $\mu$ g/ml, 45°C, 1 hr.) and solubilization in 1% sodium dodecyl sulfate (SDS). Following phenol  
25 extraction, the preparations were treated with 50 $\mu$ g/ml RNase (60°C, 30 min.), phenol/chloroform extracted, and ethanol precipitated. Standard procedures were used for enzymatic reactions and for isolation of  $\lambda$  phage DNA (Molecular Cloning Maniatis et al. Cold Springs  
30 Harbor Lab., 1982). Chlamydial DNA from serovar L<sub>2</sub> (150 $\mu$ g) was partially digested with DNase I as previously described (Ruther et al. (1982) Proc. Natl. Acad. Sci. USA 79:6852). Digested DNA was fractionated in a 1.25% agarose gel, and 500-2000 base pair  
35 fractions were collected on Whatman DE-81 paper and eluted as previously described (Dretzen et al. (1981) Annals of Biochem. 112:295). After treatment with DNA

polymerase I, the DNA was methylated with EcoRI methylase, and 2 $\mu$ g of this preparation were ligated to phosphorylated EcoRI linkers with T4 ligase. These fragments were then cleaved with EcoRI endonuclease and fractionated on a Sepharose<sup>®</sup> G-150 column. Chlamydial DNA fractions were pooled and ethanol precipitated, and 20ng of the chlamydial DNA were ligated to 1 $\mu$ g of EcoRI cleaved  $\lambda$ gt11. The EcoRI site is located within the  $\beta$ -galactosidase gene under the regulatory control of the  $\beta$ -galactosidase promoter and terminator. The ligated DNA was packaged into phage according to the manufacturer's instructions. Phage were plated and amplified in E. coli Y1088, and approximately  $2 \times 10^5$  recombinant phages were obtained.

#### 15 Screening of Recombinant Phages

E. coli Y1090 was infected with recombinant phage preparations that resulted in approximately  $10^4$  plaque forming units (PFU) per 150mm plate. Plates were initially incubated at 42°C until small plaques became visible (approx. 5 hrs.). Plates were then overlaid with IPTG saturated nitrocellulose disks and incubated an additional 2 hrs. at 37°C. The nitrocellulose disks were carefully removed from the plates, rinsed in phosphate buffered saline (PBS) (pH 7.4) to remove any residual agar, and blocked in PBS containing 5% bovine serum albumin (BSA) for 60 min. at 37°C to prevent subsequent nonspecific adsorption of protein. The disks were incubated with monoclonal antibodies (1:1000 dilution in PBS containing 0.05% Tween<sup>®</sup>-20) for 2 hr., at room temperature or overnight at 4°C. The disks were washed for 1 hr. with 6 changes of PBS-Tween<sup>®</sup> and incubated with peroxidase-conjugated, anti-mouse antibody (1:2000) for 1 hr. at room temperature, followed by a 1 hr. incubation with peroxidase anti-peroxidase (PAP) (1:2000 dilution). The disks were then washed with 6 changes of PBS-Tween<sup>®</sup> followed by 2 changes of PBS. The immune reactions

were detected by adding 0.5mg/ml of 4-chloro-1-naphthol and 0.001%  $H_2O_2$  in PBS and agitated for 5-15 min. Plaques showing positive reactions were selected, plated at low densities, and reassayed with antibody.

- 5 This process was repeated until all plaques were reactive.

#### Analysis of Proteins by SDS-PAGE and Immunoblotting

- Lysogens were produced from selected  $\lambda$ gt11 recombinants by infecting E. coli Y1089 as previously described by Young and Davis (1983) Proc. Natl. Acad. Sci. USA 80:1194. Lysates from induced recombinant lysogens were prepared, and 20 $\mu$ g aliquots were electrophoresed on 7.5% or 10% SDS-polyacrylamide gels (SDS-PAGE) according to Laemmli (1970) Nature 227:680.
- 15 The proteins in some gels were stained with Coomassie brilliant blue, while those from other gels were electrophoretically transferred to nitrocellulose for immunoblotting, as described by Towbin (1979) Proc. Natl. Acad. Sci. USA 76:4350-4354. Following
- 20 electrophoretic transfer, nitrocellulose sheets were blocked in 5% BSA and probed with either a 1:1000 dilution of rabbit polyvalent anti-C. trachomatis antiserum or mouse ascites containing high titered monoclonal antibody. Immune reactions were detected as
- 25 described above for the screening of recombinant plaques, except that the PAP step was omitted. Prestained molecular weight standards were: myosin (200,000), phosphorylase B (92,500), BSA (68,000), ovalbumin (43,000), chymotrypsinogen (25,700),
- 30 lactoglobulin (18,400), and cytochrome C (12,300) (Bethesda Research Laboratories).

#### Characterization of $\lambda$ gt11/L2/33 Insert DNA

- $\lambda$ gt11/L2/33 insert DNA was obtained from EcoRI digests of the recombinant phage and separated on agarose gels. For dot blot hybridization,  $^{32}P$ -labelled
- 35 insert DNA was reacted with lysates of C. trachomatis serovars A/G-17/OT, B/TW-5/OT, Ba/AP-2/OT, C/TW-3/OT,



D/UW-3/Cx, E/UW-5/Cx, F/UW-6/Cx, G/UW-57, H/UW-43/Cx, I/UW-12/Ur, J/UW-36/Cx, K/UW-53/Cx, L<sub>1</sub>/440/Bu, L<sub>2</sub>/434/Bu, L<sub>3</sub>/404/Bu, C. psittaci strain Mn, and HeLa 229 host cells. Lysates were prepared from

5 approximately 10 $\mu$ g of each chlamydial strain by proteinase K digestion (1mg/ml in 10mM Tris, pH 8.5, and 1mM EDTA) for 1.5 hr. at 37°C. Samples were made to 0.2N NaOH, heated to 100°C for 5 min., and placed on ice. The NaOH was neutralized with one volume of cold

10 0.2M acetic acid, followed by 0.5 volume of cold 20XSSC. The samples were filtered through nitrocellulose sheets and the sheets were washed with 6XSSC, air dried and baked 3 hr. at 80°C. The sheets were probed with <sup>32</sup>P-labelled  $\lambda$ gt11/L2/33 insert at

15 65°C by standard procedures (Molecular Cloning supra.) Southern blots of BamHI-digested C. trachomatis DNA and endonuclease restriction mapping of the  $\lambda$ gt11/L2/33 insert were performed by standard procedures (Molecular Cloning supra.).

20 Insertion of Chlamydial DNA into  $\lambda$  1059

A library of chlamydial genomic DNA was produced in the bacteriophage lambda 1059 system, which cloning system was described by Karn et al. (1980) Proc. Natl. Acad. Sci. USA 77:5172-5176. C.

25 trachomatis L<sub>2</sub> DNA was randomized by partial digestion with endonuclease restriction enzyme Sau3A or cleaved with BamHI and ligated to BamHI digested vector. Ligated DNA was packaged in vitro, as described by Sternberg et al. (1977) Gene 1:255-280, and plated in

30 E. coli Q359 for screening as described in Karn et al. (1980) supra. Phages were plated in E. coli Q359 at densities of approximately  $3 \times 10^3$  plaque forming units per 150mm plate. The plates were overlaid with nitrocellulose disks and the disks containing plaque

35 adsorbed DNA were air dried and baked 3hr at 80°C. The disks were probed with <sup>32</sup>P labelled  $\lambda$ gt11/L2/33 insert DNA at 60°C by standard procedures (Molecular Cloning,

supra). Several plaques that produced strong signals were picked and reassayed as above until all plaques from a clone were uniformly reactive. DNA was isolated from the selected phage recombinants by standard procedures (Molecular Cloning, supra). Two clones were mapped by endonuclease restriction analysis and Southern blotting by standard procedures (Molecular Cloning, supra). Both  $\lambda$ 1059 recombinants had more than one BamHI insert, however, bands with identical gel mobilities were identified which were not shared with bands from the vector, and some of these bands hybridized to the insert DNA probes in Southern blots. This process permitted mapping of contiguous endonuclease restriction sites that flanked the location of the homolog to the  $\lambda$ gt11/L2/33 insert. The map obtained by endonuclease restriction analyses was verified by generating subclones of specific fragments in a plasmid vector (pUC 18), and predicted cross-hybridizations between these clones and with the  $\lambda$ 1059 recombinants were observed in Southern blots. Fragments that included the putative coding region and flanking regions were used for DNA sequencing.

### RESULTS

#### Detection of Chlamydial Antigens

DNA obtained from C. trachomatis serovar L<sub>2</sub> was partially digested with DNase I and inserted into the bacteriophage vector  $\lambda$ gt11. The resulting plaques were transferred to nitrocellulose for the direct detection of C. trachomatis-specific antigens. Polyvalent anti-L<sub>2</sub> rabbit serum detected seven plaques that produced strong immune reactions from among the  $2 \times 10^4$  recombinant plaques assayed. The positive plaques were replated at low densities and screened with polyvalent antiserum. After plaque purification, the seven recombinants were tested with a pool of monoclonal antibodies. The monoclonal antibody pool consisted of

four antibodies (2C1, 2G1, 2H2, AE11) that each bind a mutually exclusive MOMP determinant (Stephens et al. (1982) supra.) On of the clones, designated

λgt11/L2/33, reacted with the pool of antibodies, while

5 the other six recombinant clones did not.

Subsequently, λgt11/L2/33 was tested with each of over 15 monoclonal antibodies representing species-,

subspecies-, and type-specific anti-chlamydial reaction patterns. The specificities of the antibodies and

10 their reaction pattern with λgt/11/L2/33 are presented in Table 1. The reaction pattern demonstrated that

λgt11/L2/33 was producing a polypeptide that displays species-, subspecies-, and type-specific epitopes of the chlamydial MOMP. The lack of reaction of

15 λgt11/L2/33 to antibodies not reactive with the L<sub>2</sub> serovar was expected since the recombinant was derived from serovar L<sub>2</sub> DNA. Two antibodies (AE11 and 3H10)

that do react with native L<sub>2</sub> MOMP did not react with the polypeptide expressed by λgt11/L2/33 using this

20 plaque assay. The two antibodies, however, gave positive reactions with λgt11/L2/33 expressed polypeptide in immunoblotting.

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Table 1

	Monoclonal	Serovar	Reaction with
	<u>Antibody No.</u>	<u>Specificities</u>	<u>λgt11/L2/33</u>
	2C1, IH8	all serovars	+
5	AE11	all serovars except C	-*
	3H10	A,B,D,E,F,G,H,K,L1,L2,L3	-*
	KG5	B,D,E,F,G,H,K,L1,L2,L3	+
	DA10	B,D,E,G,F,L1,L2,L3	+
	2G3	B,D,E,K,L1,L2,L3	+
10	2G1	B,F,G,H,K,L2,L3	+
	3H1, 2IIE3	B,D,E,L1,L2	+
	JC8	B,D,G,F,L2	+
	FE10	E,G,F,L2	+
	JG1	B,D,E,L2	+
15	2H2, 2H5	L2	+
	1B7, DD1	B	-
	2B1	C,J	-
	FC2	F	-
	JG9	D	-

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\* Positive reaction obtained by immunoblotting.

#### Analysis of Recombinant Fusion Polypeptides

25 E. coli lysogens were prepared for each of the positive λgt11 clones to provide a source of fusion polypeptides for analysis. Lysates obtained from induced lysogens were assessed by Coomassie blue stained PAGE gels and by immunoblotting of the proteins that were electrophoretically transferred from PAGE

30 gels to nitrocellulose. The molecular weights of these fusion proteins were estimated to range from 132,000 to 146,000.

Immunoblot analysis of PAGE gels using polyvalent rabbit anti-L<sub>2</sub> revealed that each of the

35 seven clones produced strong reactions in the plaque assay. The λgt11/L2/33 product stained most intensely, while the products from two other recombinants stained

very faintly. Immunoblot analysis was also performed with the monoclonal antibodies. Of the seven recombinants, only  $\lambda$ gt11/L2/33 reacted with monoclonal antibodies as expected from the results on the plaque assays with these same antibodies. The monoclonal antibodies that recognized species-specific and subspecies-specific determinants on L<sub>2</sub> chlamydial MOMP reacted strongly with the polypeptide produced by  $\lambda$ gt11/L2/33, while the L<sub>2</sub> type-specific monoclonal antibodies produced negative or equivocal reactions.

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Characterization of  $\lambda$ gt11/L2/33 Insert DNA

DNA from the  $\lambda$ gt11/L2/33 recombinant was isolated, labelled with <sup>32</sup>P, and used to probe dot blots of each of the 15 C. trachomatis serovars, the Mn strain of C. psittaci, and HeLa 229 host cells. Reactions were detected with all chlamydiae but not with HeLa 229 host cell DNA. Furthermore, Southern blots of BamHI digests of C. trachomatis DNA obtained from serovars L<sub>2</sub>, B, and C revealed one fragment in each preparation which reacted with <sup>32</sup>P-labelled insert DNA from  $\lambda$ gt11/L2/33. The molecular weight of this fragment varied slightly between serovars but was approximately 9.4 kb.

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Preparations of  $\lambda$ gt11/L2/33 insert DNA were obtained from EcoRI digests and separated on agarose gels. The insert was estimated to be about 1.1kb in length with restriction sites for HaeII, HaeIII, HhaI, and XhoI. Restriction sites for AccI, BamHI, BclI, BstEII, EcoRI, EcoRV, PstI, PvuI, SstI, and SstII were not detected.

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The approximately 1.1 kb insert DNA was sequenced by standard techniques, and the sequence is set forth in Appendix A.

Sequencing of  $\lambda$  1059 Inserts

Lambda 1059 recombinants having 9.2 to 9.8 kb inserts that were shown to be homologous with  $\lambda$ gt11/L2/33 by Southern analysis were used for

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endonuclease restriction mapping, and additional  
Southern analyses. Two contiguous fragments  
(BamHI/EcoRI and EcoRI/EcoRI) were identified, and  
these contain sufficient base pairs to encode for the  
5 L<sub>2</sub> MOMP gene product. These fragments were cloned into  
M13 for DNA sequencing. The sequence data for a 9.2 kb  
fragment (designated L2 B9-F DNA) are set forth in  
Appendix B.

The sequence includes an untranslated region  
10 comprising 1287 bases, followed by a 66 base region  
encoding a 22 amino acid leader sequence. Coding for  
the MOMP begins at base 67 (amino acid 23) and extends  
through base number 1182 (amino acid 394). The  
molecular weight for the MOMP including the leader is  
15 calculated to be 42,557 daltons.

The N-terminus of the MOMP was located on the  
basis of the 25 amino acid N-terminus reported by Nano  
et al. (1985) supra. Differences in the sequences of  
the N-terminus reported by Nano et al. and that  
20 reported herein are found at amino acid residues 32,  
44, and 45, as numbered in Appendix B. These  
differences may result from differences among the  
isolates or mistakes in amino acid sequencing.

The sequence set forth in Appendix A  
25 corresponds to amino acids 247 through the 3'-terminus  
in Appendix B, with certain deviations. Bases 36-38 in  
Appendix A are AGA, corresponding to amino acids  
GlyGlu, while bases 773-775 in Appendix B are TGT,  
corresponding to amino acids GlyVal. These deviations  
30 are underlined in both Appendices. The DNA sequence  
corresponding to amino acids 305 through 394 in  
Appendix B has several deviations from Appendix A which  
result in a different reading frame for the sequence of  
Appendix B. Base numbers 174, 181, and 186 in Appendix  
35 A were not detected in the  $\lambda$  1059 clones. Base number  
35 in Appendix A is a T, while the corresponding base  
in Appendix B (in amino acid 357) is a C. Finally, a G  
is inserted in amino acid 358 and a G is inserted in

amino acid 374 in the sequence of Appendix B. In both Appendices A and B, bases which are inserted or changed relative to the other Appendix are boxed, while deleted bases are indicated by an arrow. Both the DNA and amino acid sequences of Appendix B are believed to be correct.

According to the subject invention, novel recombinant DNA constructs are provided for the expression of a polypeptide having immunological activity corresponding to that of a naturally-occurring major outer membrane protein of Chlamydia trachomatis. Such polypeptides may find use as reagents in the detection of Chlamydia trachomatis or antibodies to Chlamydia trachomatis, and as vaccines against infection by Chlamydia trachomatis in susceptible hosts.

Although the foregoing invention has been described in some detail by way of illustration and example for purposes of clarity of understanding, it will be obvious that certain changes and modifications may be practiced within the scope of the appended claims.

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1 GluPheProLeuAspLeuLysAlaGlyThrAspGlyGluThrGlyThrLysAspAlaLeu  
GAATTCCTCTTTGATCTTAAAGCAAGAACAGATGGAGAGACAGGAACCTAAAGGATGCCTCT  
CTTAAGGGAGAACTAGAAATTCGTCTCTTGTCTACCTCTCTGTCTTGTCTTACCTACGGAGA  
1 ecor1, 7 mnl1, 13 mbo1 sau3a, 47 dde1, 51 fok1, 56 mnl1.

61 IleAspTyrHisGluTrpGlnAlaSerLeuAlaLeuSerTyrArgLeuAsnMetPheThr  
ATTGATTACCATGAATGGCAAGCAAGTTTAGCTCTCTCTTACAGACTGAATATGTTTCACT  
TAACTAATGGTACTTACCGTTTCGTTCAAATCGAGAGAGAATGTCTGACTTATACAAGTGA  
90 alu1, 108 xmn1,

121 ProTyrIleGlyValLysTrpSerArgAlaSerPheAspAlaAspThrIleArgTyrCys  
CCCTACATTGGAGTTAAATGGTCTCGAGCAAGTTTTGATGCAGACACGATTCTGTTATTGC  
GGGATGTAACCTCAATTTACCAGAGCTCGTTCAAACCTACGTCTGTGCTAAGGATAACG  
143 avai xho1, 144 taq1, 168 hinf1, 180 mnl1,

1 1 LeuSerProLysSerAlaThrThrValPheAspValThrThrLeuAsnProThrIleAla  
GTCAGCCCGAAGTCAGCTACAACGTCTTTGATGTTACCACTCTGAACCCCAACTATTGCT  
GAGTGGGCTTCAGTCGATGTTGACAGAACTACAATGGTGAGACTTGGGTTGATAACGA  
181 dde1, 195 alu1,

241 GlyAlaGlyAspValLysAlaSerAlaGluGlyGlnLeuGlyAspThrMetGlnIleVal  
GGAGCTGGCGATGTGAAAGCTAGCGCAGAGGGTCAGCTCGGAGATACCATGCCAAATCGTT  
CCTCGACCGCTACACTTTTCGATCGCGTCTCCAGTCGAGCCTCTATGGTACGTTTAGCAA  
243 alu1, 258 alu1, 263 hha1, 268 mnl1, 275 alu1,

301 SerLeuGlnLeuAsnLysMetLysSerArgLysPheSerValLeuGlnAM  
TCCTTGCAATTGAACAAGATGAAATCTAGAAAATTTTCGGTATTGCAGTAGGAACAACCTA  
AGGAACGTTAACTTGTCTACTTTAGATCTTTTAAAGCCATAACGTCATCCTTGTGAT  
325 xba1, 329 xmn1,

361 TTGTGGATGCAGACAAATACGCATTACAGTTGAGACTCGCTTGATCGATGAGAGAGCTGC  
AACACCTACGTCTGTTTATGCGTAATGTCAACTCTGAGCGAACTAGCTACTCTCTCGACG  
365 fok1, 394 hinf1, 403 mbo1 sau3a, 404 cla1, 405 taq1, 415  
alu1, 416 bbv fnu4h1,

421 TCACGTAAATGCACAATTCCGCTTCTAATTAATTGTATAATTTTGTAAACTTTGGCAAG  
AGTGCATTTACGTGTTAAGGGCAAGATTAAATTAACATATTTAAACAATTTGAAACCGTTC

481 TTTATCTTTGTTAATAACGTTAATAACACTATCCGTGTTTCTGGGCTCGACTTCGGTCCG  
AAATAGAAACAATTATTGCAATTATTGTGATAGGCACAAAGACCCGAGCTGAAGCCAGCC  
523 ban2 hsiJ11 sdu1, 527 taq1, 539 ban1, 540 asu1 ava2,

541 GTCCAGTTTTTTTTTGCAAAAATTTTTTTTCTTACTTTTCGATCTCCCTCCTATCTCTCTTA  
CAGGTCAAAAAAACGTTTTTAAAAAAGAATGAAAGCTAGAGGGAGGATAGAGAGAAT  
577 taq1, 579 mbo1 sau3a, 585 mnl1,

601 CAACAAAATCTAAAATTTCTCTAAAAGAAGATTGCATAAAAGGCCTCTTTCCAGTACTAT  
GTTGTTTTAGATTTTAAAGAGATTTTCTTCTAACGTATTTTCCGGAGAAAGGTCATGATA  
627 mbo1, 641 hae1 stu1, 642 hae111, 644 mnl1, 653 sca1, 65  
4 rsal,

661 ATCGGTCTACTTGAGCGCGCCCGTAGCTCAATGGTAGAGCTGTAGCCCTTCCAAGCTACCG  
TAGCCAGATGAACTCGCGCGGGCATCGAGTTACCATCTCGACATCGGAAGGTTTCGATGGC  
665 acc1, 675 hha1, 676 tac1, 677 hha1, 685 alu1, 698 alu1,  
713 alu1, 718 hpa11,



APPENDIX B

GGATCCTCACCCTCTTCATAAGCAGCAATGCATTCTCTTAGGTTTCCTAACTCCCTCGT  
AATTTTTCTATGTTCTTCTGCGCTAATAGGTCGACATACCCCAACAAATCAGCTACTG  
TGGGCCCTTCAGGATAATGTCTGCGAACCAGGATTCCACATGCAATCCTGGCCAATCC  
TTCTCCAGCATCTTGAGCCTTAAAAAGTAGGCTCAGATACATTAGGTCCTGAAGGATGT  
AAGGTACAGACCCCTAATACAGAAGCTTTGGCATGGATCGTATCTTCAACGAAATCACGG  
TCCATATGCAGCTCTTGAGCCAAAAAATCTTGCAAATTTTTTAATGTAATCCTTACGAA  
CAGGAACAAGCCTCTTATTCCCTTGCTCATCTGTATGCCAAACACGGCTCGGAATATCA  
CGTATCGCTCGATAAGAGATCCCCACATTATATGTAAACGTTTTCTGCCAGCGTTT  
ACCAAAACGATCACATACTCCTGCGCGATCACAAATGCTCAGGAACACTTCTACGCTGAG  
GACGATAAGCTTCTTCTTCTTTTCTCATGCTGACAAACAGCAACATGCCATATACGG  
AAAGTGATGATAGACAAAGCAATAATAAATCCACAAAGTAGTCTGTAGCCTTTTCTGG  
CACAGAAAGTGGGGTGGCTCGTTTTCTTTTCATATATGGTTAGTTTAATCTGTTTTTAT  
TGGTCGACCGTTTAAAAACACTTTCTTTGTAGTAATAAAGGATTTCTATCAAAACAA  
ATTCTTAGATTTTCTTACAAAATCTCTCTTTTCTTTTAGCCAAACCCCATCTTCCA  
GCTATTCCAAACACAAAAATCTTAGGTTTGGAAATTAACAACCTCATAAAAATGAACT  
GTTTTGTAAATTAACTCAAAACCTCTCATCTCAACAAATCAACATATTGCCAATATGGC  
TTTTGCTCTCGGTTTTCAGAGCGATTTTTTTTGGCAAAAGCCAAACATAAACATAAAA  
AGATATACAAAAATGGCTCTCTGCTTTATCGCTAAATCAGGAGGCGCTTAAGGGCTTCT  
TCCTGGGACCAACGTTTTTCTTATCTTCTTTACGAGAAATAAGGAAATTTTGTATGGCT  
CGAGCATTGAACGACATGTTTCGATTAAAGGCTGCTTTTACTTGCAAGACATTCTCAG  
GCCATTAATTGCTACAGGACATCTTGTCTGGCTTTAACTAGGAGGCACTGGCGCCAGAA

AAAGATAGCGAGGCACAAAGAGAGCCTAATTATACAATTTAGAGGTAAGA															1 Met ATG	Lys AAA
Lys	Leu	Leu	Lys	Ser	Val	Leu	Val	Phe	Ala	Ala	Leu	Ser	Ser	Ala		
AAA	CTC	TTG	AAA	TGG	GTA	TTA	GTG	TTT	GCC	GCT	TTG	AGT	TCT	GCT		
															10	
Ser	Ser	Leu	Gln	Ala	Leu	Pro	Val	Gly	Asn	Pro	Ala	Glu	Pro	Ser		
TCC	TCC	TTG	CAA	GCT	CTG	CCT	GTG	GGG	AAT	CCT	GCT	GAA	CTA	AGC		
															20	
Leu	Met	Ile	Asp	Gly	Ile	Leu	Trp	Glu	Gly	Phe	Gly	Gly	Asp	Pro		
CTT	ATG	ATC	GAC	GGA	ATT	CTA	TGG	GAA	GCT	TTC	GGC	GGA	CAT	CCT		
															30	
Cys	Asp	Pro	Cys	Thr	Thr	Trp	Cys	Asp	Ala	Ile	Ser	Met	Arg	Met		
TGC	CAT	CCT	TGC	ACC	ACT	TGG	TGT	GAC	GCT	ATC	AGC	ATG	CGT	ATG		
															40	
Gly	Tyr	Tyr	Gly	Asp	Phe	Val	Phe	Asp	Arg	Val	Leu	Gln	Thr	Asp		
GGT	TAC	TAT	GCT	GAC	TTT	GTT	TTC	GAC	GCT	GTT	TTG	CAA	ACA	GAT		
															50	
Val	Asn	Lys	Glu	Phe	Gln	Met	Gly	Ala	Lys	Pro	Thr	Thr	Ala	Thr		
GTG	AAT	AAA	GAA	TTC	CAA	ATG	GCT	GCC	AGG	CCT	ACA	ACT	GCT	ACA		
															60	
															70	
															80	
															90	

100  
 y Asn Ala Ala Ala Pro Ser Thr Cys Thr Ala Arg Glu Asn Pro  
 GGC AAT GCT GCA GCT CCA TCC ACT TGT ACA GCA AGA GAG AAT CCT  
 110  
 Ala Tyr Gly Arg His Met Gln Asp Ala Glu Met Phe Thr Asn Ala  
 GCT TAC GGC CGA CAT ATG CAG GAT GCT GAG ATG TTT ACA AAT GCT  
 120  
 Ala Tyr Met Ala Leu Asn Ile Trp Asp Arg Phe Asp Val Phe Cys  
 GCT TAC ATG GCA TTG AAT ATT TCG GAT CGT TTT GAT GTA TTC TGT...  
 130  
 Thr Leu Gly Ala Thr Ser Gly Tyr Leu Lys Gly Asn Ser Ala Ser  
 ACA TTA GGA CCC ACC AGT GGA TAT CTT AAA GGA AAT TCA GCA TCT  
 140  
 Phe Asn Leu Val Gly Leu Phe Gly Asp Asn Glu Asn His Ala Thr  
 TTC AAC TTA GTT GGC TTA TTC GGA GAT AAT GAG AAC CAT GCT ACA  
 150  
 Val Ser Asp Ser Lys Leu Val Pro Asn Met Ser Leu Asp Gln Ser  
 GTT TCA GAT AGT AAG CTT GTA CCA AAT ATG AGC TTA GAT CAA TCT  
 160  
 Val Val Glu Leu Tyr Thr Asp Thr Thr Phe Ala Trp Ser Ala Gly  
 GTT GTT GAG TTG TAT ACA GAT ACT ACT TTT GCT TGG AGT GCT GGA  
 170  
 Ala Arg Ala Ala Leu Trp Glu Cys Gly Cys Ala Thr Leu Gly Ala  
 GCT CGT GCA GCT TTG TGG GAA TGT GGA TGC GCG ACT TTA GGC GCT  
 180  
 Ser Phe Gln Tyr Ala Gln Ser Lys Pro Lys Val Glu Glu Leu Asn  
 TCT TTC CAA TAC GCT CAA TCC AAG CCT AAA GTC GAA GAA TTA AAC  
 190  
 Val Leu Cys Asn Ala Ala Glu Phe Thr Ile Asn Lys Pro Lys Gly  
 GTT CTC TGT AAC GCA GCT GAG TTT ACT ATC AAT AAG CCT AAA GGA  
 200  
 Tyr Val Gly Gln Glu Phe Pro Leu Asp Leu Lys Ala Gly Thr Asp  
 TAT GTA GGG CAA GAA TTC CCT CTT GAT CTT AAA GCA GGA ACA GAT  
 210  
 Gly Val Thr Gly Thr Lys Asp Ala Ser Ile Asp Tyr His Glu Trp  
 GGT GTG ACA GGA ACT AAG GAT GCC TCT ATT GAT TAC CAT CAA TGG  
 220  
 Gln Ala Ser Leu Ala Leu Ser Tyr Arg Leu Asn Met Phe Thr Pro  
 CAA GCA AGT TTA GCT CTC TCT TAC AGA CTG AAT ATG TTC ACT CCC  
 230  
 Tyr Ile Gly Val Lys Trp Ser Arg Ala Ser Phe Asp Ala Asp Thr  
 TAC ATT GGA GTT AAA TGG TCT CGA GCA AGT TTT GAT GCA GAC ACG  
 240  
 Ile Arg Ile Ala Gln Pro Lys Ser Ala Thr Thr Val Phe Asp Val  
 ATT CGT ATT GCT CAG CCG AAG TCA GCT ACA ACT GTC TTT GAT GTT  
 250  
 Thr Thr Leu Asn Pro Thr Ile Ala Gly Ala Gly Asp Val Lys Ala  
 ACC ACT CTG AAC CCA ACT ATT GCT GGA GCT GGC GAT GTG AAA GCT  
 260  
 Ser Ala Glu Gly Gln Leu Gly Asp Thr Met Gln Ile Val Ser Leu  
 AGC GCA GAG GGT CAG CTC GGA GAT ACC ATG CAA ATC GTT TCC TTG  
 270  
 Gln Leu Asn Lys Met Lys Ser Arg Lys Ser Cys Gly Ile Ala Val  
 CAA TTG AAC AAG ATG AAA TCT AGA AAA TTT TTT GGT ATT GCA GTA  
 280  
 Gly Thr Thr Ile Val Asp Ala Asp Lys Tyr Ala Val Thr Val Glu  
 GGA ACA ACT ATT GTG GAT CCA GAC AAA TAC GCA GTT ACA GTT GAG

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Thc Arg <sup>380</sup> Leu Ile Asp Glu Arg Ala Ala His Val Asn <sup>390</sup> Ala Gln Phe  
ACT CGC TTG ATC GAT GAG AGA GCT GCT CAC GTA AAT GCA CAA TTC

Arg <sup>394</sup> Phe OC  
CGC TTC TAA TTAATTGTATAATTTTGTAAACTTTGGCAAGTTTATCTTTGTAAATG  
ACGTTAATAACACTATCCGTGTTTCTGGGCTCGACTTCCGGTCCGGTCCAGTTTTTTTTTGC  
AAAAATTTTTTTTCTTACTTTTCGATCTCCCTCCTATCTCTCTTACAACAAATCTAAAT  
TTCTCTAAAAGAAGATTGCATAAAAGGCCTCTTTCCAGTACTATATCGGTCTACTTGAGG  
GCCCCCGTAGCTCAATGGTAGAGCTGTAGCCCTTCCAAGCTACCGGTGTCAAGTTCGATTCT  
GATCGGGCTCTTTTTTACTCCTGTATGACTCCCAAGTCTGAAATCTGAGCGTCTCTCAOA  
TGCCTTGTTAACACATAAAAAGAGGAACAAAGCTTGGAACTTTCTTGCAAACTCACTT-1A  
AAAGAACTATTAGAATCCGGGGCACATTTTGGACACCAGACAGTCCGCTGGAAATCCCAAG  
ATGAAGCCTTTTATTTTTGAAGAAAAAATGGCCTTTACATCATCGACTTGGCTAAAAC  
TTAGGTCAGTTGAAAAAGGCTGTTTCTTGCAATCAAAAAACTATCGATCAAGAGAGGTCT  
ATTTTTGTTTGTGGAACAAAAAACAAGCAAAACAGATCATTAGAGAAGCTGCTATCGA  
ATGTGGCGAATTC

WHAT IS CLAIMED IS:

1. A polypeptide composition capable of eliciting the production of antibodies to a major outer membrane protein (MOMP) of Chlamydia trachomatis, said polypeptide composition including non-interfering amounts of a microorganism other than Chlamydia trachomatis.
2. A polypeptide as in Claim 1, which polypeptide is produced by introducing into the unicellular host a DNA construct including a MOMP polynucleotide.
3. A polypeptide as in Claim 2, wherein the MOMP polynucleotide is at least 27 base pairs in length.
4. A polypeptide as in Claim 2, wherein the MOMP polynucleotide encodes for substantially the entire length of a major outer membrane protein.
5. A polypeptide as in Claim 2, wherein the MOMP polynucleotide is fused to a second structural gene in proper reading frame therewith and is under the regulatory control of the regulatory system of the second structural gene.
6. A polypeptide as in Claim 5, wherein the second structural gene codes for an immunogen.
7. A polypeptide as in Claim 5, wherein the second structural gene is a  $\beta$ -galactosidase gene.
8. A polypeptide as in Claim 1, wherein the unicellular host is Escherichia coli.

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9. A polypeptide as in Claim 1, wherein the immunological activity is cross-reactive with Chlamydia trachomatis serovar L<sub>2</sub>.

5           10. A DNA construct comprising a MOMP polynucleotide coding for at least a portion of the amino acid sequence of a major outer membrane protein of Chlamydia trachomatis, said DNA fragment being flanked at its 5'-end with transcriptional and  
10           translational initiation regulatory sequences and at the 3'-end with transcriptional and translational termination regulatory sequences, wherein at least one of said transcriptional or translational regulatory sequences is not derived from Chlamydia trachomatis.

15           11. A DNA construct as in Claim 10, wherein the MOMP polynucleotide is at least 27 base pairs in length and encodes for an epitopic site of the MOMP.

20           12. A DNA construct as in Claim 10, wherein the MOMP polynucleotide encodes for substantially the entire length of a major outer membrane protein.

25           13. A DNA construct as in Claim 10, wherein the MOMP polynucleotide is fused to a second structural gene in proper reading frame therewith and is under the regulatory control of the regulatory system of the second structural gene.

30           14. A DNA construct as in Claim 13, wherein the second structural gene codes for an immunogen.

35           15. A DNA construct as in Claim 13, wherein the second structural gene is a  $\beta$ -galactosidase gene.

16. A DNA construct as in Claim 6, further comprising a replication system.

17. A DNA construct as in Claim 16, wherein the replication system is a prokaryotic replication system.

5 18. A DNA construct as in Claim 17, wherein the prokaryotic replication system is recognized by Escherichia coli.

10 19. A DNA construct as in Claim 18, wherein the prokaryotic replication system is derived from  $\lambda$  phage.

20. A polynucleotide encoding a polypeptide having immunological activity cross-reactive with a  
15 major outer membrane protein of Chlamydia trachomatis, said polynucleotide being substantially free of natural flanking regions.

20 21. A polynucleotide as in Claim 20, wherein the major outer membrane protein is from Chlamydia trachomatis serovar L<sub>2</sub>.

25 22. A polynucleotide as in Claim 20, having a nucleotide sequence substantially as set forth in Appendix B hereto, or a portion thereof including at least 12 bases.

30 23. An immunoassay for the detection of Chlamydia trachomatis or antibodies to Chlamydia trachomatis in a biological sample, said immunoassay characterized by the use as reagents of the polypeptides of Claim 1, where the polypeptides may be labelled or unlabelled.

35 24. A polynucleotide probe for detecting the presence of Chlamydia trachomatis in a biological sample, said probe comprising a single stranded

polynucleotide fragment capable of binding a DNA or RNA sequence characteristic of one or more species of Chlamydia trachomatis, and a detectable label bound to the single-stranded polynucleotide.

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25. A polynucleotide probe as in claim 24, wherein the sequence of the single stranded polynucleotide fragment is substantially homologous or complementary to at least 12 contiguous bases as set forth in Appendix B hereto.

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